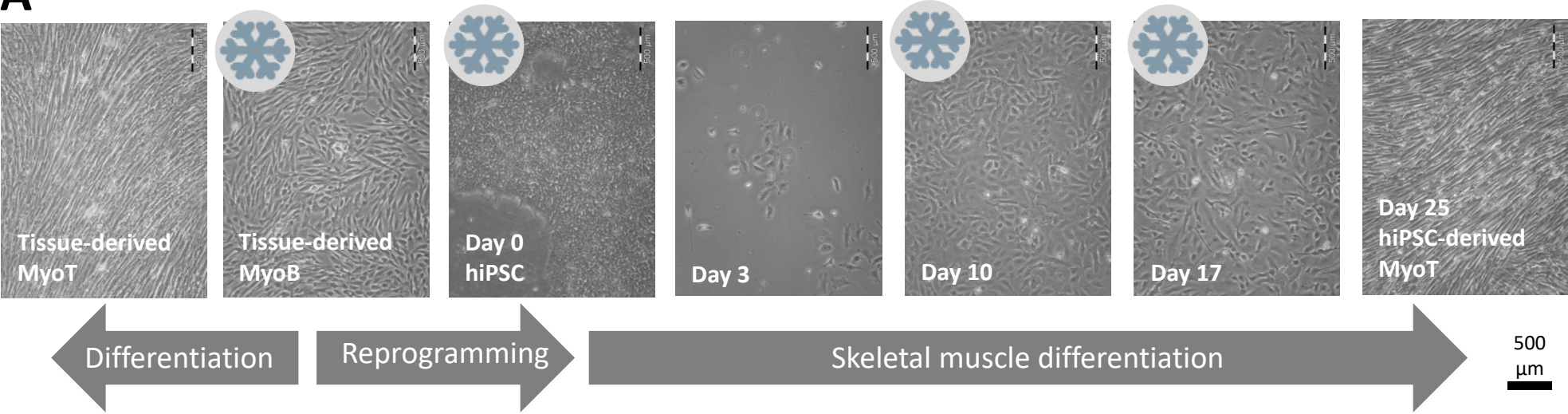


Figure S1

A



B

Target Name	Cell state	Phenotype	Ct mean	Ct SEM
<i>UBC</i> (reference gene)	D0	Healthy	23.7	0.1
		DMD	24.5	0.3
	D3	Healthy	24.9	0.3
		DMD	24.8	0.3
	D10	Healthy	24.9	0.2
		DMD	25.5	0.4
	D17	Healthy	25.7	0.6
		DMD	26.0	0.7
	D25	Healthy	25.8	0.4
		DMD	27.1	1.4
<i>Dp71-40</i>	D0	Healthy	25.2	0.3
		DMD	26.1	0.0
	D3	Healthy	25.2	0.2
		DMD	26.4	0.3
	D10	Healthy	24.5	0.7
		DMD	24.5	0.2
<i>Dp116</i>	D17	Healthy	27.2	0.5
		DMD	26.5	0.4
	D25	Healthy	26.6	0.8
		DMD	25.9	1.9
<i>Dp140</i>	D3	Healthy	35.9	0.3
		DMD	33.6	2.3
	D10	Healthy	33.6	2.3
		DMD	35.0	0.5
<i>Dp260</i>	D17	Healthy	37.9	2.6
		DMD	35.6	0.4
	D25	Healthy	34.2	1.2
		DMD	35.3	2.5
<i>Dp412e</i>	D3	Healthy	32.1	1.6
		DMD	33.8	0.7
	D10	Healthy	32.7	0.7
		DMD	36.5	0.2
	D25	Healthy	32.7	2.8
<i>Dp427c</i>	D0	Healthy	34.4	0.7
		DMD	36.5	0.2
	D3	Healthy	31.2	0.5
		DMD	31.6	1.0
	D10	Healthy	29.3	1.4
		DMD	30.0	2.3
	D17	Healthy	30.0	1.0
		DMD	30.3	0.6
	D25	Healthy	25.3	1.2
		DMD	26.4	0.3
<i>Dp427m</i>	D0	Healthy	22.8	0.4
		DMD	25.1	2.5
	D3	Healthy	37.2	1.7
		DMD	35.8	2.4
	D10	Healthy	33.3	1.9
		DMD	32.7	2.1
	D17	Healthy	33.3	0.8
		DMD	34.1	0.2
<i>Dp427m</i>	D3	Healthy	38.2	3.8
		DMD	33.5	0.9
	D10	Healthy	34.8	0.4
		DMD	30.1	1.0
	D17	Healthy	30.8	0.4
		DMD	30.1	0.3
	D25	Healthy	30.1	0.3
		DMD	30.0	0.3
<i>Dp260</i>	D3	Healthy	35.2	0.9
		DMD	34.7	0.9
	D10	Healthy	28.5	1.2
		DMD	29.4	0.8
	D17	Healthy	22.5	1.1
		DMD	23.6	0.1
	D25	Healthy	20.4	0.4
		DMD	21.9	1.4

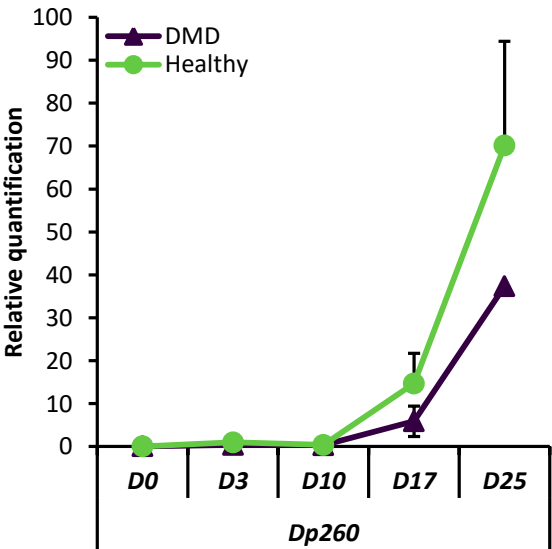
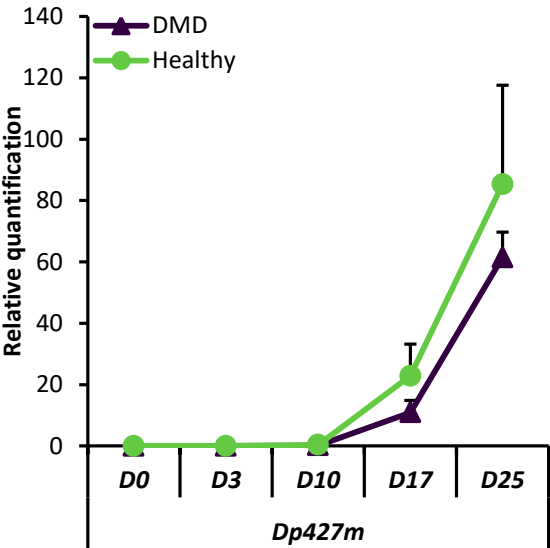
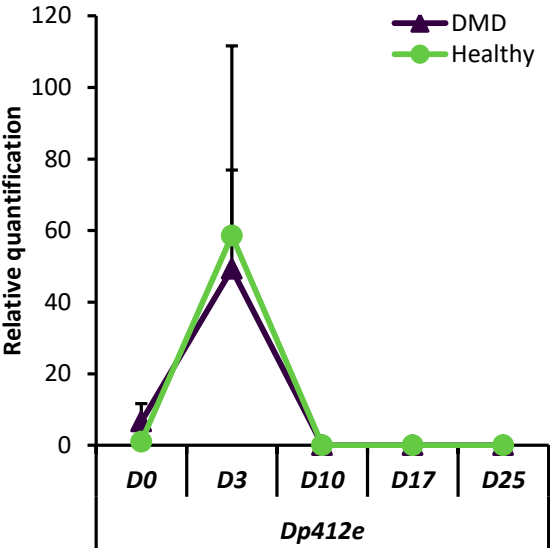


Figure S2

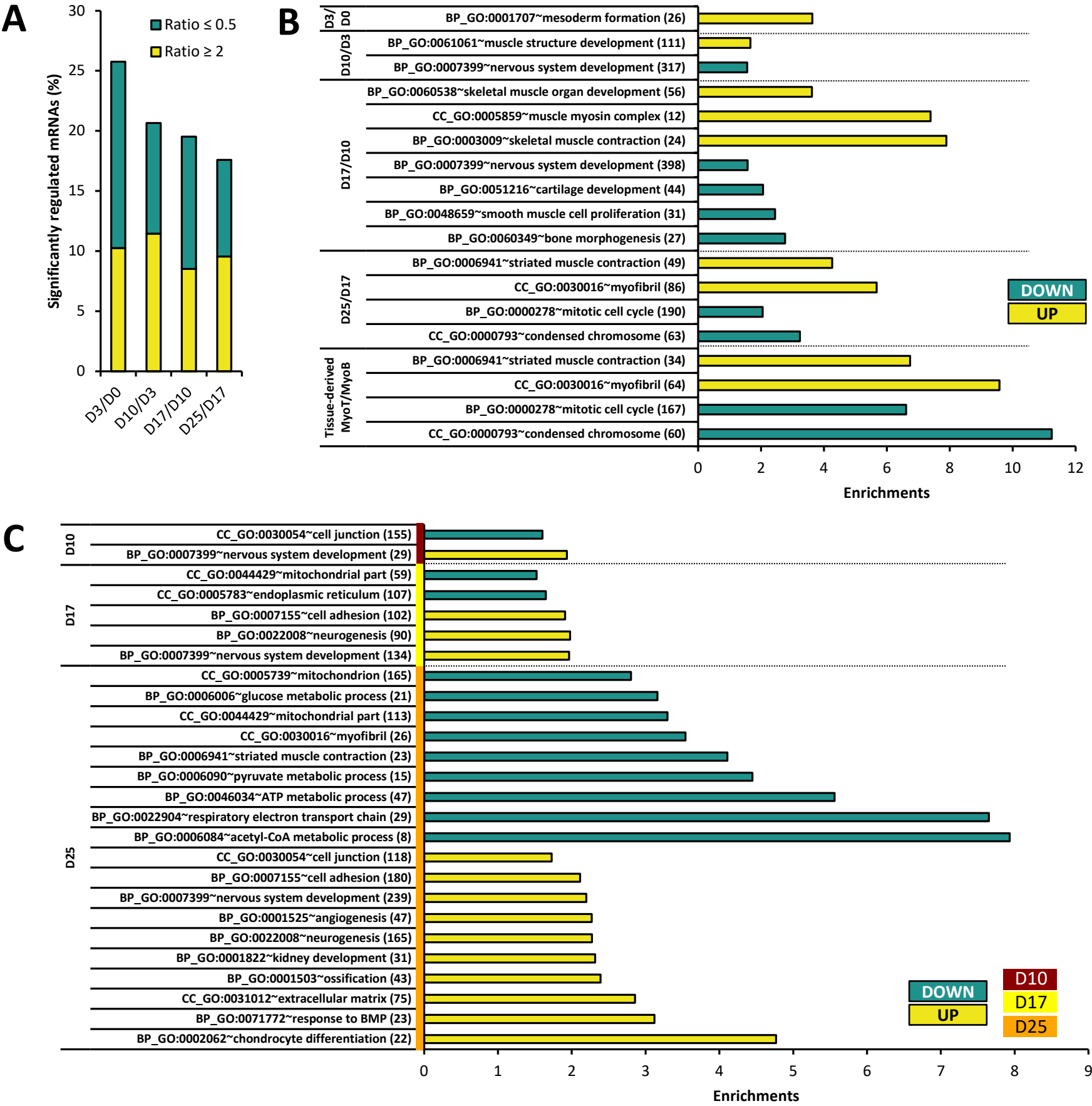
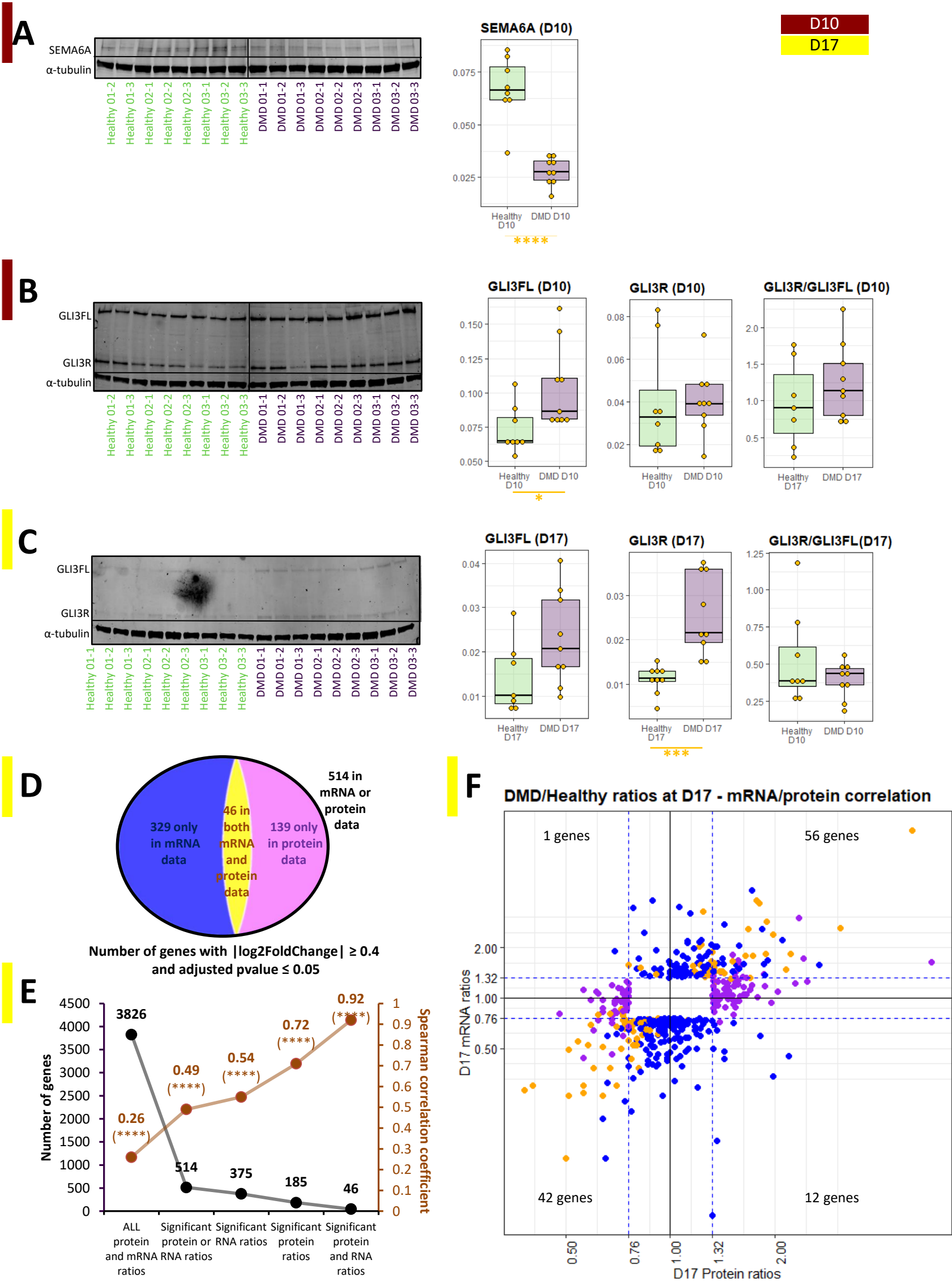
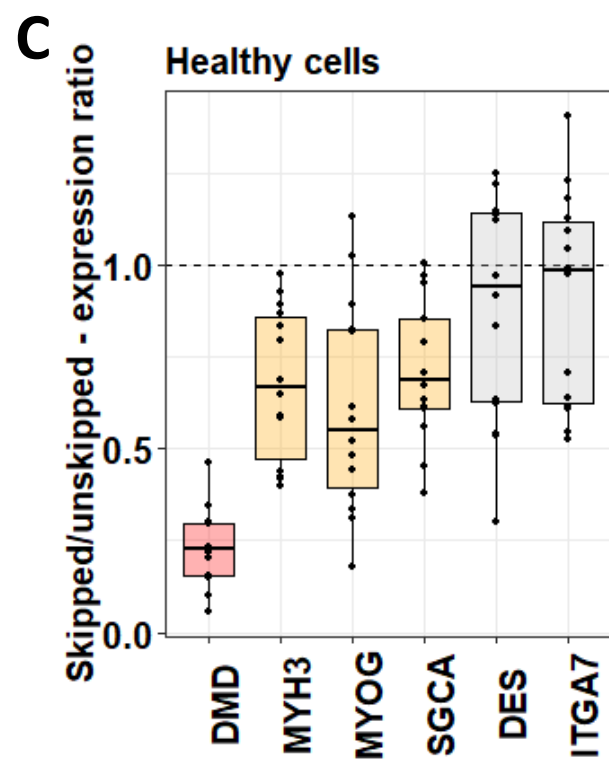
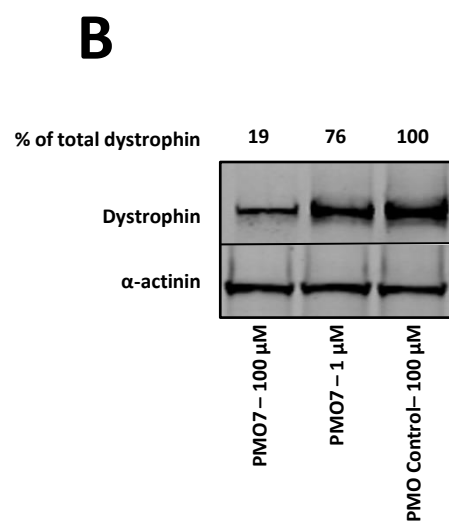
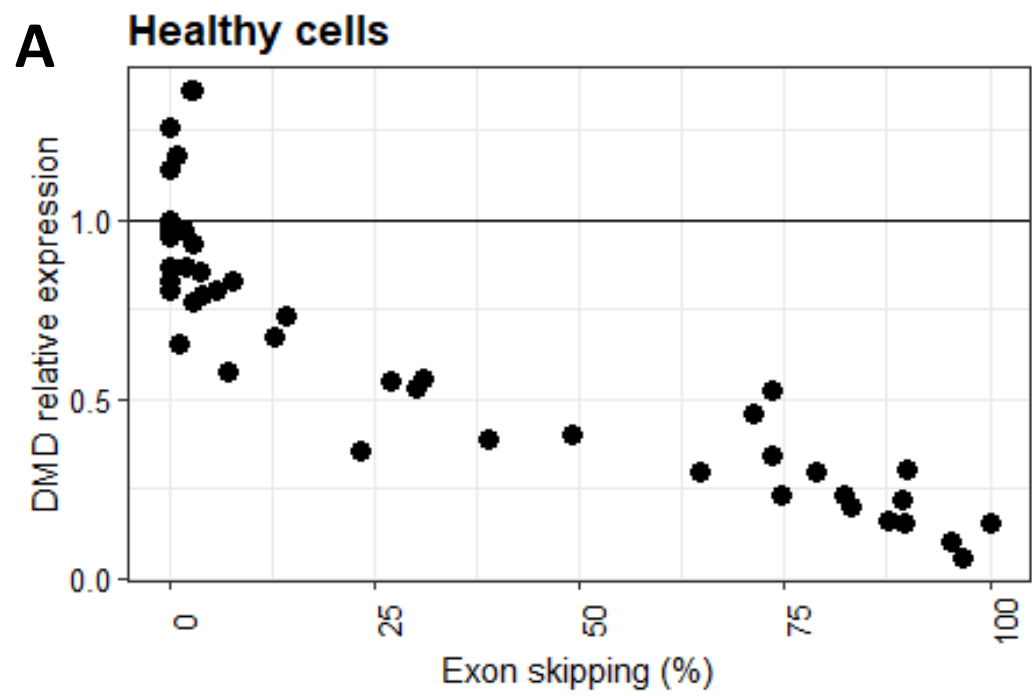


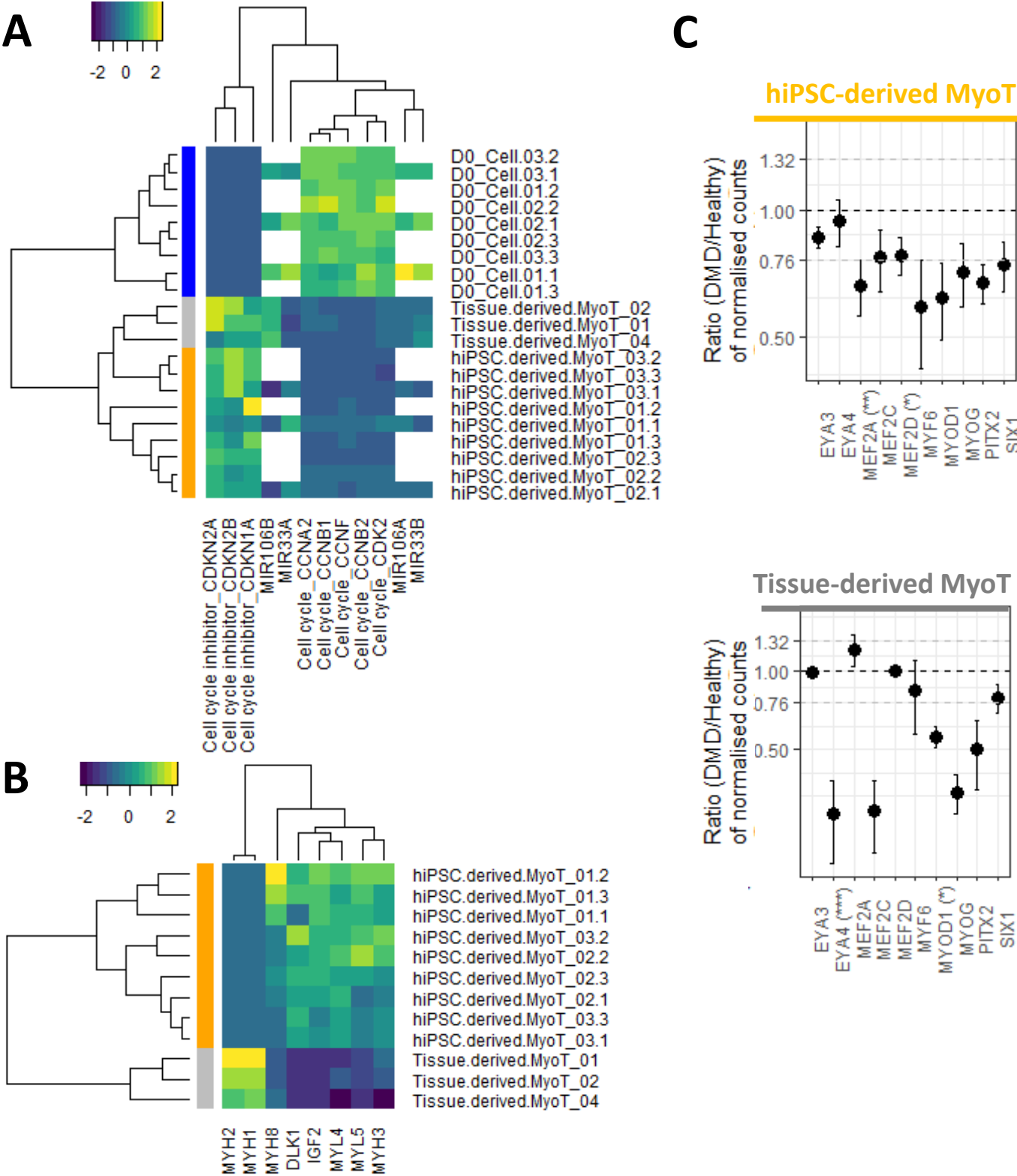
Figure S3



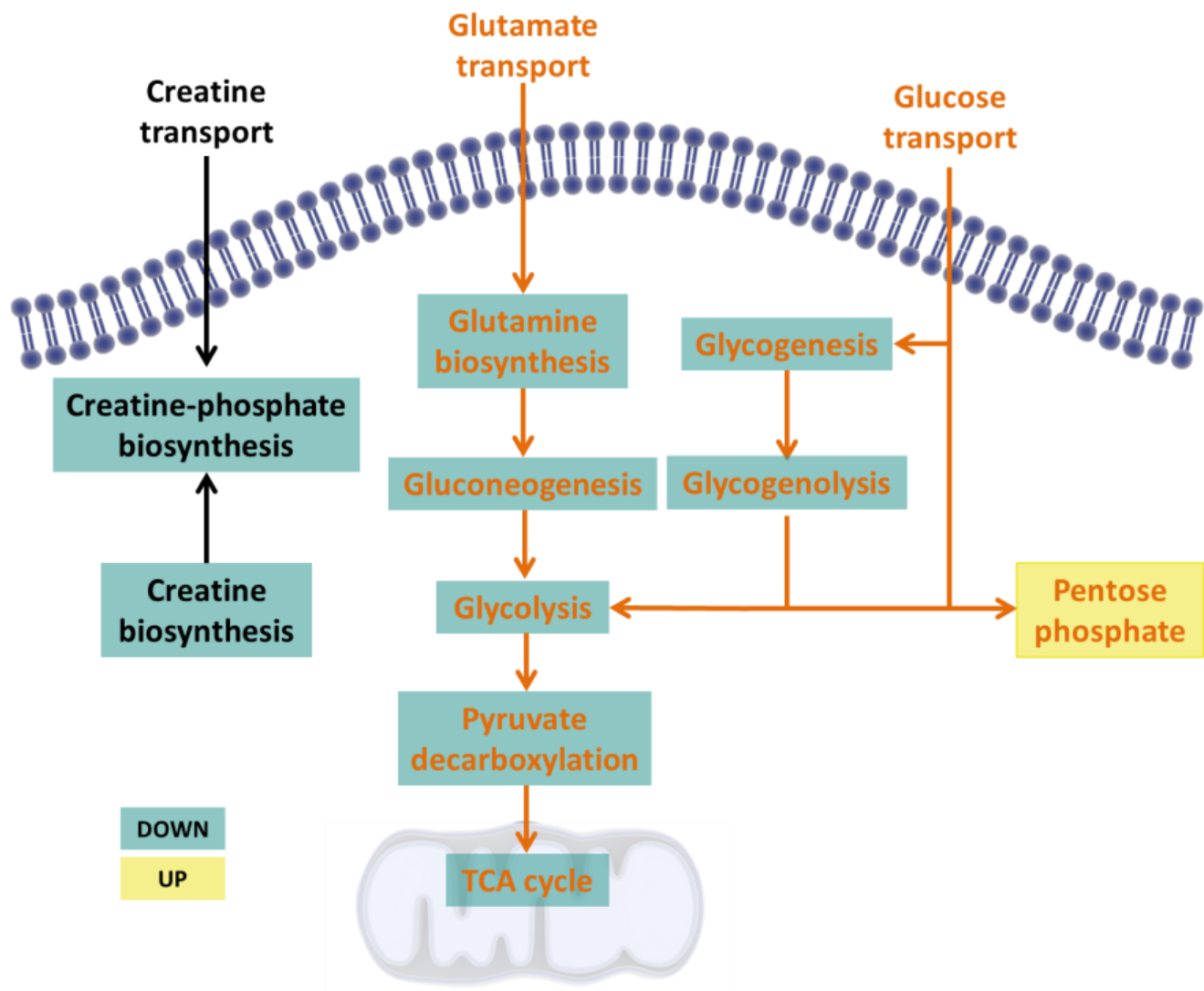


P value	****	****	****	ns	ns
Spearman r	0.56	0.64	0.5	0.15	0.062
Number of analysed pairs	59	59	56	59	59
	Group 1			Group 2	

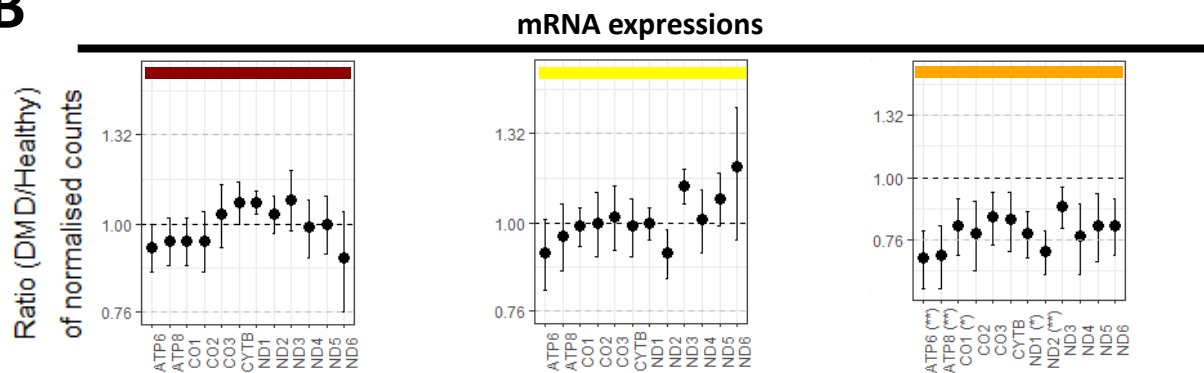
Figure S4



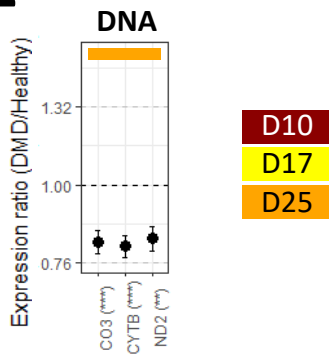
A



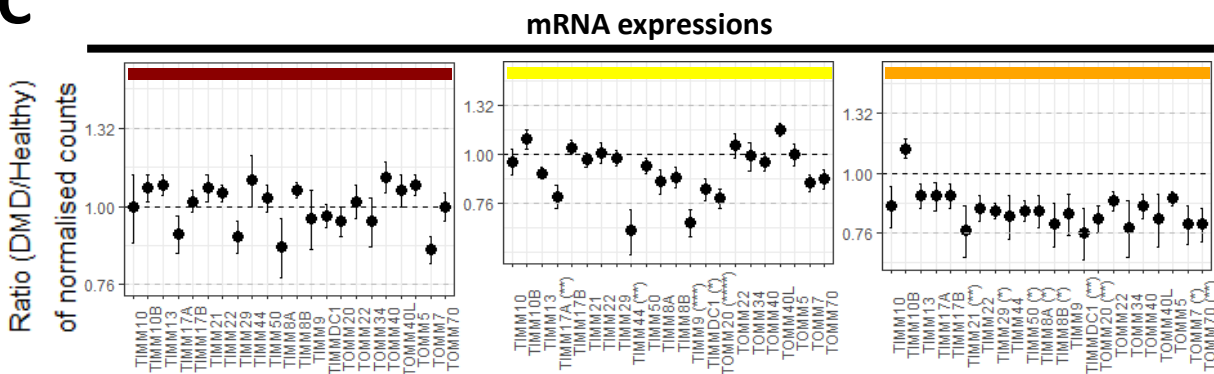
B



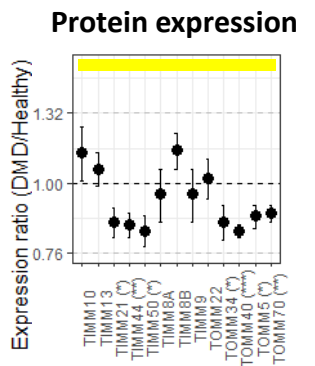
E



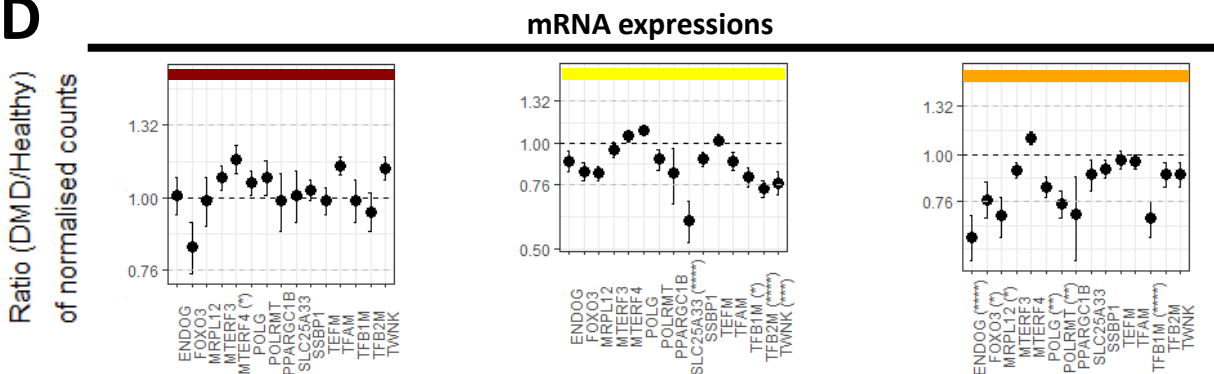
C



F



D



G

